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- => file biosis caplus caba agricola
- => s cry2? and processing
- L1 18 CRY2? AND PROCESSING
- => duplicate remove 11
- L2 10 DUPLICATE REMOVE L1 (8 DUPLICATES REMOVED)
- => d ti 1-10
- ANSWER 1 OF 10 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation of TI Interaction of two Bacillus thuringiens is delta-endotoxins with the
- TI Interaction of two Bacillus thuringiensis delta-endotoxins with the digestive system of Lygus hesperus.
- L2 ANSWER 2 OF 10 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Contrary to other non-photic cues, acute melatonin injection does not induce immediate changes of clock gene mRNA expression in the rat suprachiasmatic nuclei
- L2 ANSWER 3 OF 10 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on STN
- TI Circadian profile and photic regulation of clock genes in the suprachiasmatic nucleus of a diurnal mammal Arvicanthis ansorgei.
- L2 ANSWER 4 OF 10 CABA COPYRIGHT 2005 CABI on STN
- TI Evidence for multiple mechanisms of resistance to CrylAc and Cry2A toxins from Bacillus thuringiensis in Heliothis virescens.
- L2 ANSWER 5 OF 10 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on STN
- TI Physiological and molecular detection of crystalliferous Bacillus thuringiensis strains from habitats in the South Central United States.
- L2 ANSWER 6 OF 10 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Posttranslational mechanisms regulate the mammalian circadian clock
- L2 ANSWER 7 OF 10 CABA COPYRIGHT 2005 CABI on STN
- TI [Application of a PCR-based method for the detection of genetically modified soyabean and maize in animal feeds].

  Soia e mais geneticamente modificati: applicazione di una metodica PCR in alimenti ad uso zootecnico.
- L2 ANSWER 8 OF 10 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on STN
- TI Production of chymotrypsin-resistant Bacillus thuringiensis Cry2Aa1 delta-endotoxin by protein engineering.
- L2 ANSWER 9 OF 10 CABA COPYRIGHT 2005 CABI on STN
- TI Interaction of Bacillus thuringiensis [delta]-endotoxins with midgut brush border membrane vesicles of Helicoverpa armigera.
- L2 ANSWER 10 OF 10 CAPLUS COPYRIGHT 2005 ACS on STN
- TI The sequence of a 36 kb segment on the left arm of yeast chromosome X identifies 24 open reading frames including NUC1, PRP21 (SPP91), CDC6, CRY2, the gene for S24, a homolog to the aconitase gene ACO1 and two homologues to chromosome III genes
- => d bib abs 9 8 4 1
- L2 ANSWER 9 OF 10 CABA COPYRIGHT 2005 CABI on STN
- AN 2000:57120 CABA
- DN 20001107788
- TI Interaction of Bacillus thuringiensis [delta]-endotoxins with midgut brush border membrane vesicles of Helicoverpa armigera
- AU Shahid Karim; Riazuddin, S.; Dean, D. H.

- CS National Centre of Excellence in Molecular Biology, University of the Punjab, Canal Bank Road, Lahore-53700, Pakistan.
- SO Journal of Asia-Pacific Entomology, (1999) Vol. 2, No. 2, pp. 153-162. 40 ref.
- DT Journal
- LA English
- ED Entered STN: 20000511 Last Updated on STN: 20000511
- AB The pesticidal activity of different Bacillus thuringiensis (Bt) [delta]-endotoxins, CrylAa, CrylAb, CrylAc and Cry2A, was studied against Helicoverpa armigera infesting cotton crops worldwide. The CrylAc toxin was the most potent. All selected Bt toxins were stable to in vitro processing by midgut juice of H. armigera. Saturation and competition binding experiments were performed with iodine-125 labelled proteins and brush border membrane vesicles prepared from the midgut of H. armigera. The results showed saturable, specific and high affinity binding of all toxins except for Cry2A. Both toxins were bound with low binding affinity but with high binding site concentration. Heterologous competition experiments showed that CrylAa, CrylAb and CrylAc recognized or shared the same binding site which was different to that of Cry2A. The data suggested that the development of multiple toxin systems in transgenic plants with toxin pyramiding, which recognize different binding sites, may be useful in deployment strategies to
- L2 ANSWER 8 OF 10 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on STN

decrease the rate of pest adaptation to Bt toxins in transgenic plants.

- AN 1999:492087 BIOSIS
- DN PREV199900492087
- TI Production of chymotrypsin-resistant Bacillus thuringiensis Cry2Aa1 delta-endotoxin by protein engineering.
- AU Audtho, Mongkon; Valaitis, Algimantas P.; Alzate, Oscar; Dean, Donald H. [Reprint author]
- CS Department of Biochemistry, Ohio State University, 484 West 12th Ave., Columbus, OH, 43210-1292, USA
- SO Applied and Environmental Microbiology, (Oct., 1999) Vol. 65, No. 10, pp. 4601-4605. print.

  CODEN: AEMIDF. ISSN: 0099-2240.
- DT Article
- LA English
- ED Entered STN: 16 Nov 1999 Last Updated on STN: 16 Nov 1999
- AΒ Cleavage of the Cry2Aa1 protoxin (molecular mass, 63 kDa) from Bacillus thuringiensis by midgut juice of gypsy moth (Lymantria dispar) larvae resulted in two major protein fragments: a 58-kDa fragment which was highly toxic to the insect and a 49-kDa fragment which was not toxic. In the midgut juice, the protoxin was processed into a 58-kDa toxin within 1 min, but after digestion for 1 h, the 58-kDa fragment was further cleaved within domain I, resulting in the protease-resistant 49-kDa fragment. Both the 58-kDa and nontoxic 49-kDa fragments were also found in vivo when 125I-labeled toxin was fed to the insects. N-terminal sequencing revealed that the protease cleavage sites are at the C termini of Tyr49 and Leu144 for the active fragment and the smaller fragment, respectively. To prevent the production of the nontoxic fragment during midgut processing, five mutant proteins were constructed by replacing Leul44 of the toxin with Asp (L144D), Ala (L144A), Gly (L144G), His (L144H), or Val (L144V) by using a pair of complementary mutagenic oligonucleotides in PCR. All of the mutant proteins were highly resistant to the midgut proteases and chymotrypsin. Digestion of the mutant proteins by insect midgut extract and chymotrypsin produced only the active 58-kDa fragment, except that L144H was partially cleaved at residue 144.
- L2 ANSWER 4 OF 10 CABA COPYRIGHT 2005 CABI on STN
- AN 2003:165607 CABA
- DN 20033141535
- TI Evidence for multiple mechanisms of resistance to CrylAc and Cry2A toxins from Bacillus thuringiensis in Heliothis virescens

- AU Jurat-Fuentes, J. L.; Gould, F. L.; Adang, M. J.
- CS Department of Entomology, University of Georgia, Athens, GA 30602, USA.
- SO Resistant Pest Management Newsletter, (2003) Vol. 12, No. 2, pp. 42-44. 14 ref.

Publisher: Center for Integrated Plant Systems. East Lansing

- CY United States
- DT Journal
- LA English
- ED Entered STN: 20031003
  - Last Updated on STN: 20031003
- AB Toxin-binding assays using radiolabelled CrylA toxins were conducted to study the mechanism of resistance to the bacterial toxins in H. virescens strains CXC and KCBhyb. The strains were isolated and incubated with increasing concentrations of labelled CrylA toxins to generate binding saturation curves. Results indicated the presence of at least 2 resistance mechanisms in the larvae from the KCBhyb strain, one of which would be related to Cry 1A receptor alteration and the other would be related to toxin solubilization and processing in the larval midgut. Alteration of toxin solubilization and/or processing seems to be the main mechanism of resistance in CXC.
- L2 ANSWER 1 OF 10 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on STN
- AN 2004:161171 BIOSIS
- DN PREV200400164879
- TI Interaction of two Bacillus thuringiensis delta-endotoxins with the digestive system of Lygus hesperus.
- AU Brandt, Sandra L.; Coudron, Thomas A. [Reprint Author]; Habibi, Javad; Brown, Gregory R.; Ilagan, Oliver M.; Wagner, Renee M.; Wright, Maureen K.; Backus, Elaine A.; Huesing, Joseph E.
- CS Biological Control of Insects Research Laboratory, U.S. Department of Agriculture, Agricultural Research Service, 1503 S. Providence Rd, Columbia, MO, 65203, USA coudront@missouri.edu
- SO Current Microbiology, (January 2004) Vol. 48, No. 1, pp. 1-9. print. CODEN: CUMIDD. ISSN: 0343-8651.
- DT Article
- LA English
- ED Entered STN: 24 Mar 2004 Last Updated on STN: 24 Mar 2004
- AΒ The active-toxin form of CrylAc (65 kDa) or Cry2Ab was fed to a non-susceptible insect, Lygus hesperus, in an artificial diet. Biochemical and immunocytochemical methods were used to determine the distribution of ingested toxin. The toxins did not elicit a feeding deterrent response. CrylAc and Cry2Ab were ingested; small amounts were absorbed into the hemolymph as holoproteins, but most was excreted. SDS-PAGE analysis of CrylAc and Cry2Ab incubations with salivary gland homogenate showed a small decrease in the molecular weight of the active toxins. Proteolytic **processing** of the toxins also occurred in vivo, within the digestive system of L. hesperus. Excreted CrylAc and Cry2Ab retained activity toward lepidopteran larvae. Immunocytochemical in vivo localization studies showed negligible association of CrylAc with L. hesperus tissues. In contrast, strong extracellular association of Cry2Ab was observed with L. hesperus midgut brush border microvilli and basement membrane, as well as with cellular outlines within the hemolymph and fat body.
- => s cryII? and process?
- L3 56 CRYII? AND PROCESS?
- => duplicate remove 13
- L4 28 DUPLICATE REMOVE L3 (28 DUPLICATES REMOVED)
- => d ti 1-28
- L4 ANSWER 1 OF 28 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Method for the Detection of Synthetic cry3A in Transgenic Potatoes

- L4 ANSWER 2 OF 28 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Functional analysis of two **processed** fragments of Bacillus thuringiensis CryllA toxin
- L4 ANSWER 3 OF 28 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Homologous recombination into Gram-positive bacterium for generation of expression libraries of polynucleotides
- L4 ANSWER 4 OF 28 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Methods for production and secretion of asparaginase in Bacillus subtilis
- L4 ANSWER 5 OF 28 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on STN
- TI Assessment of screening methods for the identification of genetically modified potatoes in raw materials and finished products.
- L4 ANSWER 6 OF 28 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on STN
- TI RNA processing and degradation in Bacillus subtilis.
- L4 ANSWER 7 OF 28 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on STN
- TI A detection method for recombinant DNA from genetically modified potato (NewLeaf Y(R) potato).
- L4 ANSWER 8 OF 28 CAPLUS COPYRIGHT 2005 ACS on STN
- TI PCR method for detecting recombinant DNAs from genetically modified crops and processed food
- L4 ANSWER 9 OF 28 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Bacillus thuringiensis strain fermentation **process** and insecticide application
- L4 ANSWER 10 OF 28 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Expression of multiple insecticidal genes confers broad resistance against a range of different rice pests
- L4 ANSWER 11 OF 28 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on
- TI Phase-specific optimization of multiple endotoxin-protein production with genetically engineered Bacillus thuringiensis.
- L4 ANSWER 12 OF 28 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Nucleic acid vectors for recombinant production of heterologous proteins in a Bacillus cell
- L4 ANSWER 13 OF 28 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Insect-resistant transgenic plants and methods for improving  $\delta\text{-endotoxin}$  activity against target insects
- L4 ANSWER 14 OF 28 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Transition State of the Rate-Limiting Step of Heat Denaturation of Cry3A  $\delta\textsc{-Endotoxin}$
- L4 ANSWER 15 OF 28 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on
- TI Organizational complexity of a rice transgene locus susceptible to methylation-based silencing.
- L4 ANSWER 16 OF 28 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Sporulation-incompetent strains of Bacillus thuringiensis for use in persistent  $\delta$ -endotoxin-based pesticide formulations
- L4 ANSWER 17 OF 28 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on
- TI Functional significance of loops in the receptor binding domain of Bacillus thuringiensis **CryIIIA** delta-endotoxin.
- L4 ANSWER 18 OF 28 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on
- TI Proteolytic processing of Bacillus thuringiensis CryIIIA toxin and specific binding to brush-border membrane vesicles of Leptinotarsa decemlineata (Colorado potato beetle).

- L4 ANSWER 19 OF 28 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on
- TI Transfer and transcriptional expression of coleopteran cryIIIB endotoxin gene of Bacillus thuringiensis in eggplant.
- L4 ANSWER 20 OF 28 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Safety assessment of potatoes resistant to Colorado potato beetle
- L4 ANSWER 21 OF 28 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Overproduction of encapsulated insecticidal crystal proteins in a Bacillus thuringiensis spo0A mutant
- L4 ANSWER 22 OF 28 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on
- TI Overexpression of Bacillus thuringiensis HknA, a histidine protein kinase homology, bypasses early Spo- mutations that result in **CryIIIA** overproduction.
- L4 ANSWER 23 OF 28 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on
- TI Structural and functional analysis of the promoter region involved in full expression of the **cryIIIA** toxin gene of bacillus thuringiensis.
- L4 ANSWER 24 OF 28 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on
- TI Site-directed mutations in a highly conserved region of Bacillus thuringiensis delta-endotoxin affect inhibition of short circuit current across Bombyx mori midguts.
- L4 ANSWER 25 OF 28 CABA COPYRIGHT 2005 CABI on STN
- TI The crystal [delta]-endotoxins of Bacillus thuringiensis: models for their mechanisms of action on the insect gut.
- L4 ANSWER 26 OF 28 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on
- TI Immunocytochemical localization of Bacillus thuringiensis insecticidal crystal proteins in intoxicated insects.
- L4 ANSWER 27 OF 28 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on
- TI Synthetic **cryIIIA** gene from Bacillus thuringiensis improved for high expression in plants.
- L4 ANSWER 28 OF 28 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Recovery of Bacillus thuringiensis endotoxin protein from lysed cell mixtures
- => d bib abs 2 13 18 25
- L4 ANSWER 2 OF 28 CAPLUS COPYRIGHT 2005 ACS on STN
- AN 2004:327750 CAPLUS
- DN 141:48991
- TI Functional analysis of two **processed** fragments of Bacillus thuringiensis CryllA toxin
- AU Yamagiwa, Masashi; Sakagawa, Kohei; Sakai, Hiroshi
- CS Department of Bioscience and Biotechnology, Okayama University, Okayama, 700-8530, Japan
- SO Bioscience, Biotechnology, and Biochemistry (2004), 68(3), 523-528 CODEN: BBBIEJ; ISSN: 0916-8451
- PB Japan Society for Bioscience, Biotechnology, and Agrochemistry
- DT Journal
- LA English
- AB The 70-kDa protoxin of CryllA, a dipteran-specific insecticidal protein, was processed by trypsin into 36- and 32-kDa fragments. To investigate the potent function of the two processed fragments, a GST (Glutathione-S-transferase) fusion protein of each polypeptide was constructed. While neither the 36- nor the 32-kDa fragment was toxic to Culex pipiens larvae, coexpression of the two fragments restored the insecticidal activity. Furthermore, the copptn. experiment demonstrated that the 36-kDa fragment was associated with the 32-kDa fragment. It was, therefore, shown that the coexistence of the two processed

fragments of CryllA was essential for the toxicity. The mutant of the 36-kDa fragment lacking the region from Gly257 to Arg360 bound to the 32-kDa fragment but the coexpression with the 32-kDa fragment resulted in no toxicity, suggesting that this region was involved in insecticidal activity.

RE.CNT 11 THERE ARE 11 CITED REFERENCES AVAILABLE FOR THIS RECORD ALL CITATIONS AVAILABLE IN THE RE FORMAT

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L4 ANSWER 13 OF 28 CAPLUS COPYRIGHT 2005 ACS on STN
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AN 1999:405094 CAPLUS

DN 131:55200

TI Insect-resistant transgenic plants and methods for improving  $\delta$ -endotoxin activity against target insects

IN English, Leigh; Brussock, Susan M.; Malvar, Thomas M.; Bryson, James W.;
Kulesza, Caroline A.; Walters, Frederick S.; Slatin, Stephen L.; Von
Tersch, Michael A.; Romano, Charles

PA Ecogen, Inc., USA; Monsanto Company

SO PCT Int. Appl., 512 pp.

CODEN: PIXXD2

DT Patent

LA English

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PI	WO									WO 1998-US26852						19981217			
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			KG,	KP,	KR,	KZ,	LC,	LK,	LR,	LS,	LI	Γ,	LU,	LV,	MD,	MG,	MK,	MN,	MW,
			MX,	NO,	NZ,	PL,	PT,	RO,	RU,	SD,	SE	Ξ,	SG,	SI,	SK,	SL,	ТJ,	TM,	TR,
			TT,	UA,	ŪG,	US,	US,	US,	US,	UZ,	٧N	١,	YU,	ZW,	AM,	AZ,	BY,	KG,	ΚZ,
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	US	6063597		Α	20000516			US 1997-993170					19971218						
		6077824		Α	20000620			US 1997-993775				19971218							
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		1997-993170					1997												
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		1997-993775			AI 71		1997												
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		1998-US26852																	
		1999-427770					1999												

AB Disclosed are methods for increasing the activity of Bacillus thuringiensis  $\delta$ -endotoxins against Coleopteran insect pests. The three-dimensional crystal structure of Cry3Bb  $\delta$ -endotoxin was used as the basis for protein engineering. Thirty-six mutants are created by (1) alteration of protease-sensitive sites and proteolytic processing, (2) modification of bound water and hydropathic index of amino acids, (3) manipulation of hydrogen bonds around mobile regions, (4) loop anal. and loop design around the flexible helixes and  $\beta$ -strands and  $\beta$ -sheets, (5) re-design of complex electrostatic surfaces, (6) removal of metal binding sites, (7) alteration of quaternary structure, and (8) alteration of binding to glycoproteins and to western corn rootworm brush border membranes. Also disclosed are methods for mutagenizing nucleic acid sequences encoding these polypeptides, and

increasing insect resistance in transgenic plants expressing these genes.
RE.CNT 9 THERE ARE 9 CITED REFERENCES AVAILABLE FOR THIS RECORD
ALL CITATIONS AVAILABLE IN THE RE FORMAT

- L4 ANSWER 18 OF 28 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on
- AN 1996:314786 BIOSIS
- DN PREV199699037142
- TI Proteolytic processing of Bacillus thuringiensis CryIIIA toxin and specific binding to brush-border membrane vesicles of Leptinotarsa decemlineata (Colorado potato beetle).
- AU Martinez-Ramirez, A. C. [Reprint author]; Real, M. D.
- CS Dep. de Genetica, Universitat de Valencia, Dr. Moliner 50, 46100 Burjassot, Valencia, Spain
- SO Pesticide Biochemistry and Physiology, (1996) Vol. 54, No. 2, pp. 115-122. CODEN: PCBPBS. ISSN: 0048-3575.
- DT Article
- LA English
- ED Entered STN: 11 Jul 1996 Last Updated on STN: 11 Jul 1996
- AB The mode of action of Bacillus thuringiensis insecticidal proteins in lepidopteran insects is known to involve five steps: ingestion, solubilization, protease activation, binding to midgut membrane receptors, and disruption of the intestinal membrane. Two of these steps, protease activation and binding to midgut membrane receptors, have been analyzed in the major potato pest, the coleoptera Leptinotarsa decemlineata (Colorado potato beetle). Unlike recently proposed, after treatment of the coleopteran-specific B. thuringiensis toxin CryIIIA with gut content from the Colorado potato beetle, a 42-kDa processing polypeptide has been identified. The study of binding to midgut membrane receptors has demonstrated specific and saturable binding of chymotrypsinized CryIIIA to brush-border membrane vesicles from the Colorado potato beetle. The affinity constant and the concentration of binding sites values (K-d = 37.5 + 8.6 nM, R-t = 17 + 4 pmol/mg of protein) were in the range of the ones previously estimated for low affinity binding sites in lepidopteran insects. Taking into account that CryIII4 can be proteolytically processed by the Colorado potato beetle midgut proteases, along with the fact that, in our hands, binding can be demonstrated only if the toxin is chymotrypsin processed, these results suggest that the mode of action of the coleopteran-specific B. thuringiensis toxin CryIIIA is probably the same as that of lepidopteran-specific toxins.
- L4 ANSWER 25 OF 28 CABA COPYRIGHT 2005 CABI on STN
- AN 93:117618 CABA
- DN · 19931181431
- TI The crystal [delta]-endotoxins of Bacillus thuringiensis: models for their mechanisms of action on the insect gut
- AU Knowles, B. H.; Dow, J. A. T.
- CS Department of Zoology, University of Cambridge, Downing Street, Cambridge, CB2 3EJ, UK.
- SO BioEssays, (1993) Vol. 15, No. 7, pp. 469-476. 49 ref. ISSN: 0265-9247
- DT Journal
- LA English
- ED Entered STN: 19941101
  - Last Updated on STN: 19941101
- AB A model of the effects on insect gut of [delta]-endotoxins from Bacillus thuringiensis is presented, based on a recently elucidated structure for CryIIIA. The processes of solubilisation and proteolytic activation, receptor binding and formation of the toxic lesion are described. Predictions are made for the effects of the toxins on ion movement, gap junctions and osmotic events and goblet cells.

- L5 ANSWER 1 OF 3 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Functional analysis of two **processed** fragments of Bacillus thuringiensis CryllA toxin
- L5 ANSWER 2 OF 3 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Bacillus thuringiensis strain fermentation **process** and insecticide application
- L5 ANSWER 3 OF 3 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Expression of multiple insecticidal genes confers broad resistance against a range of different rice pests
- . => d bib abs 3
  - L5 ANSWER 3 OF 3 CAPLUS COPYRIGHT 2005 ACS on STN
  - AN 2001:307179 CAPLUS
  - DN 136:65624
  - TI Expression of multiple insecticidal genes confers broad resistance against a range of different rice pests
  - AU Maqbool, Shahina Bano; Riazuddin, Sheikh; Loc, Nguyen Thi; Gatehouse, Angharad M. R.; Gatehouse, John A.; Christou, Paul
  - CS Molecular Biotechnology Unit, John Innes Centre, Norwich, NR4 7UH, UK
  - SO Molecular Breeding (2001), 7(1), 85-93 CODEN: MOBRFL; ISSN: 1380-3743
  - PB Kluwer Academic Publishers
  - DT Journal
  - LA English
  - We report the simultaneous introduction of three insecticidal genes (the AΒ Bt genes crylAc and cry2A, and the snowdrop lectin gene gna) into com. important indica rice varieties M7 and Basmati 370, by particle bombardment. Transgenic plants expressed CrylAc, Cry2A and GNA at different levels, either singly or in combination at 0.03-1%, 0.01-0.5% and 0.01-2.5% of total soluble protein, resp. The transgenes showed stable transmission and expression, and R1 transgenic plants provided significant (p<0.01) protection against three of the most important insect pests of rice: rice leaf folder (Cnaphalocrocis medinalis), yellow stemborer (Scirpophaga incertulas) and brown planthopper (Nilaparvata lugens). triple transformants showed significantly (p<0.05) higher resistance to these insects than plants expressing single transgenes. Bioassays using the triple-transgenic plants showed 100% eradication of the rice leaf folder and yellow stem borer, and 25% reduction in the survival of the brown planthopper. The greatest reduction in insect survival, and the greatest reduction in plant damage, occurred in plants expressing all three transgenes. This approach maximises the utility of gene transfer technol. to introduce combinations of genes whose products disrupt different biochem. or physiol. processes in the same insect, providing a multi-mechanism defense.
  - RE.CNT 32 THERE ARE 32 CITED REFERENCES AVAILABLE FOR THIS RECORD ALL CITATIONS AVAILABLE IN THE RE FORMAT
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  - L6 82 CRY2? AND PROCESS?
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46 DUPLICATE REMOVE L6 (36 DUPLICATES REMOVED)

- => d ti 1-46
- L7 ANSWER 1 OF 46 CAPLUS COPYRIGHT 2005 ACS on STN

- TI Pulses of prolactin promoter activity depend on a noncanonical E-box that can bind the circadian proteins CLOCK and BMAL1
- L7 ANSWER 2 OF 46 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on STN
- TI Regulation of prokineticin 2 expression by light and the circadian clock
- L7 ANSWER 3 OF 46 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on STN
- TI Cryptochromes and neuronal-activity markers colocalize in the retina of migratory birds during magnetic orientation.
- L7 ANSWER 4 OF 46 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Tobacco budworm response to CryIAc and Cry2Ab toxins of Bacillus thuringiensis
- L7 ANSWER 5 OF 46 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on STN
- TI Light-response quantitative trait loci identified with composite interval and eXtreme array mapping in Arabidopsis thaliana.
- L7 ANSWER 6 OF 46 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on STN DUPLICATE 4
- TI Comparison of broiler chicken performance when fed diets containing meals of Bollgard II hybrid cotton containing Cry-X gene (Cry1Ac and Cry2Ab gene), parental line or commercial cotton.
- L7 ANSWER 7 OF 46 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on STN
- TI Retinal cryptochrome in a migratory passerine bird: a possible transducer for the avian magnetic compass.
- L7 ANSWER 8 OF 46 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on STN
- TI Enhanced expression of insecticidal crystal proteins in wild Bacillus thuringiensis strains by a heterogeneous protein p20.
- L7 ANSWER 9 OF 46 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on STN
- TI Interaction of two Bacillus thuringiensis delta-endotoxins with the digestive system of Lygus hesperus.
- L7 ANSWER 10 OF 46 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Novel Bacillus thuringiensis insecticidal proteins
- L7 ANSWER 11 OF 46 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on
- TI Blue light activates calcium-permeable channels in Arabidopsis mesophyll cells via the phototropin signaling pathway.
- L7 ANSWER 12 OF 46 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Contrary to other non-photic cues, acute melatonin injection does not induce immediate changes of clock gene mRNA expression in the rat suprachiasmatic nuclei
- L7 ANSWER 13 OF 46 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on
- TI Circadian profile and photic regulation of clock genes in the suprachiasmatic nucleus of a diurnal mammal Arvicanthis ansorgei.
- L7 ANSWER 14 OF 46 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on
- TI Clock genes and the long-term regulation of prolactin secretion: Evidence for a photoperiod/circannual timer in the pars tuberalis.
- L7 ANSWER 15 OF 46 CAPLUS COPYRIGHT 2005 ACS on STN
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- TI Further characterization of the phenotype of mCryl/mCry2-deficient mice.

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- TI Susceptibility of the pine **processionary** caterpillar
  Thaumetopoea pityocampa (Lepidoptera: Thaumetopoeidae) toward
  delta-endotoxins of Bacillus thuringiensis under laboratory conditions.
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- TI UV and blue light signalling: Pathways regulating chalcone synthase gene expression in Arabidopsis.
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- TI Expression of multiple insecticidal genes confers broad resistance against a range of different rice pests.
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- TI [Application of a PCR-based method for the detection of genetically modified soyabean and maize in animal feeds].

  Soia e mais geneticamente modificati: applicazione di una metodica PCR in alimenti ad uso zootecnico.
- L7 ANSWER 40 OF 46 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on
- TI Production of chymotrypsin-resistant Bacillus thuringiensis Cry2Aa1 delta-endotoxin by protein engineering.
- L7 ANSWER 41 OF 46 BIOSIS COPYRIGHT (c) 2005 The Thomson Corporation on
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- TI Cloning and analysis of the first cry gene from Bacillus popilliae.
- L7 ANSWER 45 OF 46 CAPLUS COPYRIGHT 2005 ACS on STN
- TI The sequence of a 36 kb segment on the left arm of yeast chromosome X identifies 24 open reading frames including NUC1, PRP21 (SPP91), CDC6, CRY2, the gene for S24, a homolog to the aconitase gene ACO1 and two homologues to chromosome III genes
- L7 ANSWER 46 OF 46 CAPLUS COPYRIGHT 2005 ACS on STN
- TI Chronopotentiometry of electrode **processes** followed by chemical reactions involving electroactive species. Reduction of hexamminechromium(III) in the presence of ethylenediaminetetraacetate
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- AN 1999:492087 BIOSIS
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- TI Production of chymotrypsin-resistant Bacillus thuringiensis Cry2Aa1 delta-endotoxin by protein engineering.
- AU Audtho, Mongkon; Valaitis, Algimantas P.; Alzate, Oscar; Dean, Donald H. [Reprint author]
- CS Department of Biochemistry, Ohio State University, 484 West 12th Ave., Columbus, OH, 43210-1292, USA
- SO Applied and Environmental Microbiology, (Oct., 1999) Vol. 65, No. 10, pp. 4601-4605. print.

  CODEN: AEMIDF. ISSN: 0099-2240.

- DT Article
- LA English
- ED Entered STN: 16 Nov 1999

Last Updated on STN: 16 Nov 1999

- Cleavage of the Cry2Aa1 protoxin (molecular mass, 63 kDa) from AΒ Bacillus thuringiensis by midgut juice of gypsy moth (Lymantria dispar) larvae resulted in two major protein fragments: a 58-kDa fragment which was highly toxic to the insect and a 49-kDa fragment which was not toxic. In the midgut juice, the protoxin was processed into a 58-kDa toxin within 1 min, but after digestion for 1 h, the 58-kDa fragment was further cleaved within domain I, resulting in the protease-resistant 49-kDa fragment. Both the 58-kDa and nontoxic 49-kDa fragments were also found in vivo when 125I-labeled toxin was fed to the insects. N-terminal sequencing revealed that the protease cleavage sites are at the C termini of Tyr49 and Leu144 for the active fragment and the smaller fragment, respectively. To prevent the production of the nontoxic fragment during midgut processing, five mutant proteins were constructed by replacing Leul44 of the toxin with Asp (L144D), Ala (L144A), Gly (L144G), His (L144H), or Val (L144V) by using a pair of complementary mutagenic oligonucleotides in PCR. All of the mutant proteins were highly resistant to the midgut proteases and chymotrypsin. Digestion of the mutant proteins by insect midgut extract and chymotrypsin produced only the active 58-kDa fragment, except that L144H was partially cleaved at residue 144.
- L7 ANSWER 17 OF 46 CABA COPYRIGHT 2005 CABI on STN
- AN 2003:165607 CABA
- DN 20033141535
- TI Evidence for multiple mechanisms of resistance to CrylAc and Cry2A toxins from Bacillus thuringiensis in Heliothis virescens
- AU Jurat-Fuentes, J. L.; Gould, F. L.; Adang, M. J.
- CS Department of Entomology, University of Georgia, Athens, GA 30602, USA.
- SO Resistant Pest Management Newsletter, (2003) Vol. 12, No. 2, pp. 42-44. 14 ref.

Publisher: Center for Integrated Plant Systems. East Lansing

- CY United States
- DT Journal
- LA English
- ED Entered STN: 20031003

Last Updated on STN: 20031003

- AB Toxin-binding assays using radiolabelled CrylA toxins were conducted to study the mechanism of resistance to the bacterial toxins in H. virescens strains CXC and KCBhyb. The strains were isolated and incubated with increasing concentrations of labelled CrylA toxins to generate binding saturation curves. Results indicated the presence of at least 2 resistance mechanisms in the larvae from the KCBhyb strain, one of which would be related to Cry 1A receptor alteration and the other would be related to toxin solubilization and processing in the larval midgut. Alteration of toxin solubilization and/or processing seems to be the main mechanism of resistance in CXC.
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